

## AMENDMENTS TO THE SPECIFICATION

Please amend the Specification to read as follows:

Replace paragraph [0074] in the Specification of the above-identified application with the following paragraph:

[0074] The wild-type summer-annual WS accession of *Arabidopsis* was transformed with constructs containing the entire *FPA* gene using *Agrobacterium*-mediated transformation. The construct was prepared using the isolated *FPA* gene (amplified from genomic *Arabidopsis* DNA using the primers ATGGCGTTATCTATGAAGCCATTCAGAGCC (SEQ ID NO:7) and TCAAGGCCCTGTCCAGCCGGAGTACC (SEQ ID NO:8)), and a 35S CaMV promoter. Successful transformants were then collected and grown in conditions to allow assessment of the *FPA* overexpression on flowering time.

Replace paragraph [0077] in the Specification of the above-identified application with the following paragraph:

[0077] Using the *FPA* polynucleotide sequence (SEQ ID NO:2), late-flowering *Arabidopsis* were generated using both antisense and cosuppression transgenes. An antisense construct was generated using the 5' coding region of *FPA* (amplified by the primers 5' AAGACTTTAAAGGAGATGTTTCAGCC (SEQ ID NO:9) and 5' CCTTTCCCATAGGTACACAACGAGC (SEQ ID NO:10)) and expressing the opposite strand under control of the CaMV 35S promoter. Primary transformants that displayed delayed flowering were selected and progeny were replanted. Upon replanting, late-flowering plants that phenocopied *fpa* mutants were isolated (i.e., flowered after producing 60 leaves).